

SEQUENCE LISTING

<110> Ono Pharmaceutical Co., Ltd.

<120> Novel Polypeptides, DNAs encoding the polypeptides, and utility of the Polypeptides

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<140> 09/380,276

<141> 1999-08-27

<150> JP 9-43143

<151> 1997-02-27

<150> PCT/JP98/00799

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<170> PatentIn version 3.0

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<223> Origin: human bone marrow stromal cell line HAS303

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Leu	Gln	Ala	Arg	Asn	Ala	Gly	Pro	Ala	Gly	Glu	Met	Val	Pro	Thr	Phe	
255						260				265						
ttc	gga	tcc	ctc	acg	cag	tcc	atc	tgt	ggc	gag	ttt	tca	gat	gcc	tgg	968
Phe	Gly	Ser	Leu	Thr	Gln	Ser	Ile	Cys	Gly	Glu	Phe	Ser	Asp	Ala	Trp	
270						275				280						
cct	ctg	atg	cag	aat	ccc	atg	ggt	ggt	gac	aac	atc	tct	ttt	tgt	gac	1016
Pro	Leu	Met	Gln	Asn	Pro	Met	Gly	Gly	Asp	Asn	Ile	Ser	Phe	Cys	Asp	
285						290				295						
tct	tat	cct	gaa	ctc	act	gga	gaa	gac	att	cat	tct	ctc	aat	cca	gaa	1064
Ser	Tyr	Pro	Glu	Leu	Thr	Gly	Glu	Asp	Ile	His	Ser	Leu	Asn	Pro	Glu	
300						305				310			315			
ctt	gaa	agc	tca	acg	tct	ttg	gat	tca	aat	agc	agt	caa	gat	ttg	gtt	1112
Leu	Glu	Ser	Ser	Thr	Ser	Leu	Asp	Ser	Asn	Ser	Ser	Gln	Asp	Leu	Val	
320						325				330						
ggt	ggg	gct	gtt	cca	gtc	cag	tct	cat	tct	gaa	aac	ttt	aca	gca	gct	1160
Gly	Gly	Ala	Val	Pro	Val	Gln	Ser	His	Ser	Glu	Asn	Phe	Thr	Ala	Ala	
335						340				345						
act	gat	tta	tct	aga	tat	aac	aac	aca	ctg	gta	gaa	tca	gca	tca	act	1208
Thr	Asp	Leu	Ser	Arg	Tyr	Asn	Asn	Thr	Leu	Val	Glu	Ser	Ala	Ser	Thr	

350	355	360	
cag gat gca cta act atg aga agc cag cta gat cag gag agt ggc gct Gln Asp Ala Leu Thr Met Arg Ser Gln Leu Asp Gln Glu Ser Gly Ala 365	370	375	1256
atc atc cac cca gcc act cag acg tcc ctc cag gta agg cag cga ctg Ile Ile His Pro Ala Thr Gln Thr Ser Leu Gln Val Arg Gln Arg Leu 380	385	390	1304
ggt tcc ctg tgaacacagc actgacttac agtagatcag aactctgttc Gly Ser Leu			1353
ccagcataag atttggggga acctgatgag tttttttt gcatcttaa taatttcttg tatgtttagt agtatgtttt aaaataaatt tcaagtattt ttttaaaaaa ctaaaaaaaa aaaaaaaaaa aaaaaaaaaa aaa			1413 1473 1496
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Leu Val Leu Leu Gly Tyr Leu Ser Cys Lys Val Thr Cys Glu Thr Gly -5	-1	1	5
Asp Cys Arg Gln Gln Glu Phe Arg Asp Arg Ser Gly Asn Cys Val Pro 10	15	20	
Cys Asn Gln Cys Gly Pro Gly Met Glu Leu Ser Lys Glu Cys Gly Phe 25	30	35	
Gly Tyr Gly Glu Asp Ala Gln Cys Val Thr Cys Arg Leu His Arg Phe 40	45	50	55
Lys Glu Asp Trp Gly Phe Gln Lys Cys Lys Pro Cys Leu Asp Cys Ala 60	65	70	
Val Val Asn Arg Phe Gln Lys Ala Asn Cys Ser Ala Thr Ser Asp Ala			

75

80

85

Ile Cys Gly Asp Cys Leu Pro Gly Phe Tyr Arg Lys Thr Lys Leu Val
 90 95 100

Gly Phe Gln Asp Met Glu Cys Val Pro Cys Gly Asp Pro Pro Pro Pro
 105 110 115

Tyr Glu Pro His Cys Ala Ser Lys Val Asn Leu Val Lys Ile Ala Ser
 120 125 130 135

Thr Ala Ser Ser Pro Arg Asp Thr Ala Leu Ala Ala Val Ile Cys Ser
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Ala Leu Ala Thr Val Leu Leu Ala Leu Leu Ile Leu Cys Val Ile Tyr
 155 160 165

Cys Lys Arg Gln Phe Met Glu Lys Lys Pro Ser Trp Ser Leu Arg Ser
 170 175 180

Gln Asp Ile Gln Tyr Asn Gly Ser Glu Leu Ser Cys Leu Asp Arg Pro
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Gln Leu His Glu Tyr Ala His Arg Ala Cys Cys Gln Cys Arg Arg Asp
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Ser Val Gln Thr Cys Gly Pro Val Arg Leu Leu Pro Ser Met Cys Cys
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Glu Glu Ala Cys Ser Pro Asn Pro Ala Thr Leu Gly Cys Gly Val His
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Ser Ala Ala Ser Leu Gln Ala Arg Asn Ala Gly Pro Ala Gly Glu Met
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Val Pro Thr Phe Phe Gly Ser Leu Thr Gln Ser Ile Cys Gly Glu Phe
 265 270 275

Ser Asp Ala Trp Pro Leu Met Gln Asn Pro Met Gly Gly Asp Asn Ile
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Ser Phe Cys Asp Ser Tyr Pro Glu Leu Thr Gly Glu Asp Ile His Ser
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Leu Asn Pro Glu Leu Glu Ser Ser Thr Ser Leu Asp Ser Asn Ser Ser
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Gln Asp Leu Val Gly Gly Ala Val Pro Val Gln Ser His Ser Glu Asn
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Ser Ala Ser Thr Gln Asp Ala Leu Thr Met Arg Ser Gln Leu Asp Gln
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28